

Draw A)

SEQUENCE LISTING

<110> SUGA, Mikiko
ASAKURA, Yoko
MORI, Yukiko
ITO, Hisao
KURAHASHI, Osamu

<120> Arginine Repressor Deficient Stain of Coryneform Bacterium and
Method for Producing L-Arginine

<130> OP1018

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<141> 2001-04-

<150> JP2000-129167

<151> 2000-04-28

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<170> PatentIn Ver. 2.0

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Pro Thr His Gly Asp Ala Thr Lys Arg Asp Leu Asn Ala Ala Val Phe	
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<213> *Brevibacterium lactofermentum*

<220>

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<211> 427

<212> PRT

<213> Brevibacterium lactofermentum

<400> 8

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Ser	Lys	Ala	Leu	Ala	Gly	Cys	His	Arg	Trp	Arg	Arg	Asp	Glu	Ala	Val
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Gly	Glu	Arg	Arg	Ile	Glu	Leu	Ala	Ile	Ala	Thr	Lys	Asn	His	Leu	Ala
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Ala	Gly	Gly	Ala	Leu	Met	Met	Phe	Val	Gly	Thr	Val	Arg	His	Asn	Arg
								100		105		110			
Ser	Gln	Ser	Phe	Ala	Gln	Val	Glu	Ala	Gly	Ile	Lys	Thr	Ala	Tyr	Ser
								115		120		125			

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 Val Glu His Thr Tyr Ser Asp Tyr Glu Val Thr Asp Ser Trp Ala Asn
 145 150 155 160
 Gly Trp His Leu His Arg Asn Met Leu Leu Phe Leu Asp Arg Pro Leu
 165 170 175
 Ser Asp Asp Glu Leu Lys Ala Phe Glu Asp Ser Met Phe Ser Arg Trp
 180 185 190
 Ser Ala Gly Val Val Lys Ala Gly Met Asp Ala Pro Leu Arg Glu His
 195 200 205
 Gly Val Lys Leu Asp Gln Val Ser Thr Trp Gly Gly Asp Ala Ala Lys
 210 215 220
 Met Ala Thr Tyr Leu Ala Lys Gly Met Ser Gln Glu Leu Thr Gly Ser
 225 230 235 240
 Ala Thr Lys Thr Ala Ser Lys Gly Ser Tyr Thr Pro Phe Gln Met Leu
 245 250 255
 Asp Met Leu Ala Asp Gln Ser Asp Ala Gly Glu Asp Met Asp Ala Val
 260 265 270
 Leu Val Ala Arg Trp Arg Glu Tyr Glu Val Gly Ser Lys Asn Leu Arg
 275 280 285
 Ser Ser Trp Ser Arg Gly Ala Lys Arg Ala Leu Gly Ile Asp Tyr Ile
 290 295 300
 Asp Ala Asp Val Arg Arg Glu Met Glu Glu Leu Tyr Lys Leu Ala
 305 310 315 320
 Gly Leu Glu Ala Pro Glu Arg Val Glu Ser Thr Arg Val Ala Val Ala
 325 330 335
 Leu Val Lys Pro Asp Asp Trp Lys Leu Ile Gln Ser Asp Phe Ala Val
 340 345 350
 Arg Gln Tyr Val Leu Asp Cys Val Asp Lys Ala Lys Asp Val Ala Ala
 355 360 365
 Ala Gln Arg Val Ala Asn Glu Val Leu Ala Ser Leu Gly Val Asp Ser
 370 375 380
 Thr Pro Cys Met Ile Val Met Asp Asp Val Asp Leu Asp Ala Val Leu
 385 390 395 400
 Pro Thr His Gly Asp Ala Thr Lys Arg Asp Leu Asn Ala Ala Val Phe
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<210> 9

<211> 30

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer for
introducing mutation to pAM330

<400> 9

aaacccgggc tacgttgtat gctttgaatc

30

<210> 10

<211> 27

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer for
introducing mutation to pAM330

<400> 10

tttgcgtttt cgttaacgtc aacaacc

27

<210> 11

<211> 28

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer for
introducing mutation to pAM330

<400> 11

ttttcccggtt agcttgccac accccgag

28

<210> 12

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer for
introducing mutation to pAM330

<400> 12
gggggtcatc tctggctgaa ttgg 24

<210> 13
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<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:primer for introducing mutation to pAM330

<400> 13
gagggtttca ccggtctgca tgcc 24

<210> 14
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:primer for introducing mutation to pAM330

<400> 14
aactcaccgc cctgcaattc aac 23

<210> 15
<211> 25
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:primer for PCR

<400> 15
gcctaccgcg gcaaagaagt ggcag 25

<210> 16
<211> 25

<212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:primer for PCR

<400> 16
 gccttgaact agggcgctt taagt

25

<210> 17
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 <213> *Brevibacterium flavum*

<220>
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 <222> (1852)..(2364)

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 tgggtccct cgcgtact ggccagccag acaagcgtga agcgttccctg ccaatgc当地 180
 gcggtgtgga gtttaccct tacggcgaca cggattactt gcgcaaaatg gtagaaacca 240
 acccaacgga tggcgctgat atcttctcg agccaatcca gggtgaaacg ggcgttgc当地 300
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 ccactttcgat tggcgatgcacttgcgttgcgttgcgttgcgttgcgttgcgttgc当地 600
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 aggttgcgttgcgttgcgttgcgttgcgttgcgttgcgttgcgttgcgttgcgttgc当地 720
 gcgacgttcgttgcgttgcgttgcgttgcgttgcgttgcgttgcgttgcgttgc当地 780
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 35 40 45 50
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 85 90 95
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 gcg atg ctg cgc acc ccg ccg gga gct gcc cag tac ctg gca agt ttc 2241
 Ala Met Leu Arg Thr Pro Pro Gly Ala Ala Gln Tyr Leu Ala Ser Phe
 115 120 125 130
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 Ile Asp Arg Val Gly Leu Lys Glu Val Val Gly Thr Ile Ala Gly Asp
 135 140 145
 gac acc gtt ttt gtt ctc gcc cgt gat ccg ctc aca ggt aaa gaa cta 2337
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<210> 18

<211> 171

<212> PRT

<213> *Brevibacterium flavum*

<400> 18

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 35 40 45
 Ile Asp Ile Thr Gln Ala Thr Leu Ser Arg Asp Leu Asp Glu Leu Gly
 50 55 60
 Ala Arg Lys Val Arg Pro Asp Gly Gly Arg Ala Tyr Tyr Ala Val Gly
 65 70 75 80
 Pro Val Asp Ser Ile Ala Arg Glu Asp Leu Arg Gly Pro Ser Glu Lys
 85 90 95
 Leu Arg Arg Met Leu Asp Glu Leu Leu Val Ser Thr Asp His Ser Gly
 100 105 110
 Asn Ile Ala Met Leu Arg Thr Pro Pro Gly Ala Ala Gln Tyr Leu Ala
 115 120 125
 Ser Phe Ile Asp Arg Val Gly Leu Lys Glu Val Val Gly Thr Ile Ala
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 145 150 155 160
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<210> 19

<211> 25

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: primer for PCR

<400> 19

cccggtttt cttctgcaac tcggg

25

<210> 20

<211> 25

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: primer for PCR

partial

<400> 20
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25

<210> 21
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:primer for PCR

<400> 21
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24

<210> 22
<211> 25
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:primer for PCR

<400> 22
gtcttacctc ggctgg~~tt~~gg ccagc

25